SEQUENCE LISTING

| 5 | (1) GENE | RAL INFORMATION: |
|----|----------|--|
| | (i) | APPLICANT: Gjerset, Ruth A. |
| | (ii) | TITLE OF INVENTION: DOWN-REGULATION OF DNA REPAIR TO ENHANCE SENSITIVITY TO P53-MEDIATED SUPPRESSION |
| 10 | | |
| | (iii) | NUMBER OF SEQUENCES: 3 |
| | (iv) | CORRESPONDENCE ADDRESS: |
| | | (A) ADDRESSEE: Arnold, White & Durkee |
| 15 | | (B) STREET: P.O\ Box 4433 |
| | | (C) CITY: Houston |
| | | (D) STATE: Texas |
| | • | (E) COUNTRY: United States of America |
| 20 | • | (F) ZIP: 77210 \ |
| 20 | (**) | COMPUTER READABLE FORM: |
| | (>) | (A) MEDIUM TYPE: Floppy disk |
| | | (B) COMPUTER: IBM RC compatible |
| | | (C) OPERATING SYSTEM: PC-DOS/MS-DOS |
| 25 | | (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 |
| | | |
| | (vi) | CURRENT APPLICATION DATA: |
| | | (A) APPLICATION NUMBER: US Unknown |
| • | | (B) FILING DATE: Condurrently Herewith |
| 30 | | (C) CLASSIFICATION: Unknown |
| | / | ADDODATES /ACTION TATEODATES |
| | (V111) | ATTORNEY/AGENT INFORMATION: (A) NAME: Highlander, \$teven L. |
| | | (B) REGISTRATION NUMBER: 37,642 |
| 35 | | (C) REFERENCE/DOCKET NUMBER: INGN:032 |
| | | (0, 100 210 210 210 210 210 210 210 210 210 |
| • | (ix) | TELECOMMUNICATION INFORMATION: |
| | | (A) TELEPHONE: (512) 418 \ 3000 |
| | | (B) TELEFAX: (512) 474-75/77 |
| 40 | | |
| | (0) | |
| | (2) INFO | RMATION FOR SEQ ID NO:1: |
| | (i) | SEQUENCE CHARACTERISTICS: |
| 45 | | (A) LENGTH: 24 base pairs |
| | | (B) TYPE: nucleic acid |
| | | (C) STRANDEDNESS: single |
| | | (D) TOPOLOGY: linear |
| 50 | (and) | CECHENCE DECORIDATION, SEC ID NO.1: |
| JU | (X1) | SEQUENCE DESCRIPTION: SEQ ID NO:1: |

| | | • | |
|--------------------|----|--|----|
| | | (2) INFORMATION FOR SEQ ID NO:2: | |
| | 5 | (i) SEQUENCE CHARACTERISTICS: | |
| | | (A) LENGTH 24 base pairs | |
| | | (B) TYPE: nucleic acid | |
| | | (C) STRANDEDNESS: single | |
| | | (D) TOPOLOGY: linear | |
| \ | 10 | | |
| - | | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: | |
| 1001 | | CARGON CACH, CHICAGOTTCA CIT. CACH | 24 |
| ~ \ \ | | GATCCACAGT CTGCCTGAGT CACT | 24 |
| 12/ | 15 | | |
| | 10 | (2) INFORMATION FOR SEQ ID NO:3: | |
| / | | (-, | |
| / | | (i) SEQUENCE CHARACTERISTICS: | |
| • | | (A) LENGTH: 24 base pairs | |
| | 20 | (B) TYPE: nucleic acid | |
| # % | | (C) STRANDEDNESS: single | |
| inal and gard gard | | (D) TOPOLOGY: limear | |
| 7. 5 | | (wi) CEOUENGE DECOREDATION, CEO ID NO.3. | |
| 1.11 1.71 | 25 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: | |
| | 23 | CCTAGAAAGC ACATGGAGAG CTAG | 24 |
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